SEQUENCE LISTING

<110> INSTITUT NATIONAL DE LA RECHERCHE AGRONOMIQUE Oulmouden, Ahmad Julien, Raymond Lafort, Marie-Piere, Leveziel, Hubert

<120> USE OF THE SILVER GENE FOR THE AUTHENTIFICATION OF THE ORIGINAL BREED OF ANIMAL POPULATIONS AND THEIR DERIVATIVE PRODUCTS

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<151> July 25, 2003

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tot otg ata tac ag gtgagatoco ogocatoctg otoccactoo tttacocott Ser Leu Ile Tyr Arg 600	7312
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gaccaccact aaccagtate cetgettite teccaatate ag g ega aga ett atg $$\operatorname{Arg}$$ Arg Leu Met $$\operatorname{605}$$	7907
aag caa ggc toa gca gtc occ ott occ oag otg oca oac ggt aga acc Lys Gln Gly Ser Ala Val Pro Leu Pro Gln Leu Pro His Gly Arg Thr 610 615 620	7955
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Pro Glu Trp Thr Glu Ser Gln Gly Pro Asp Cys Trp Arg Gly Gly His
Ile Ser Leu Lys Val Ser Asn Asp Gly Pro Thr Leu Ile Gly Ala Asn
Ala Ser Phe Ser Ile Ala Leu His Phe Pro Lys Ser Gln Lys Val Leu
Pro Asp Gly Gln Val Ile Trp Ala Asn Asn Thr Ile Ile Asn Gly Ser
Gln Val Trp Gly Gly Gln Leu Val Tyr Pro Gln Glu Pro Asp Asp Thr
Cys Ile Phe Pro Asp Gly Glu Pro Cys Pro Ser Gly Pro Leu Ser Gln
Lys Arg Cys Phe Val Tyr Val Trp Lys Thr Trp Asp Gln Tyr Trp Gln
Val Leu Gly Gly Pro Val Ser Gly Leu Ser Ile Gly Thr Asp Lys Ala
Met Leu Gly Thr Tyr Asn Met Glu Val Thr Val Tyr His Arg Arg Gly
Ser Gln Ser Tyr Val Pro Leu Ala His Ser Ser Ser Ala Phe Thr Ile
Thr Asp Gln Val Pro Phe Ser Val Ser Val Ser Gln Leu Gln Ala Leu
Asp Gly Arg Asn Lys Arg Phe Leu Arg Lys Gln Pro Leu Thr Phe Ala
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Tyr Thr Trp Asp Phe Gly Asp Ser Thr Gly Thr Leu Ile Ser Arg Ala
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Leu Thr Val Thr His Thr Tyr Leu Glu Ser Gly Pro Val Thr Ala Gln 275 280 285

Va	al.	Val 290	Leu	Gln	Ala	Ala	Ile 295	Pro	Leu	Thr	Ser	Cys 300	Gly	Ser	Ser	Pro
Va 30		Pro	Gly	Thr	Thr	Asp 310	Arg	His	Val	Thr	Thr 315	Ala	Glu	Ala	Pro	Gly 320
Th	ır	Thr	Ala	Gly	Gln 325	Val	Pro	Thr	Thr	Glu 330	Val	Met	Gly	Thr	Thr 335	Pro
G]	-У	Gln	Val	Pro 340	Thr	Ala	Glu	Ala	Pro 345	Gly	Thr	Thr	Val	Gly 350	Trp	Val
Pı	0	Thr	Thr 355	Glu	Asp	Val	Gly	Thr 360	Thr	Pro	Glu	Gln	Val 365	Ala	Thr	Ser
ΓŽ	rs	Val 370	Leu	Ser	Thr	Thr	Pro 375	Val	Glu	Met	Pro	Thr 380	Ala	Lys	Ala	Thr
G] 38		Arg	Thr	Pro	Glu	Val 390	Ser	Thr	Thr	Glu	Pro 395	Ser	Gly	Thr	Thr	Val 400
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Se	r	Thr	Pro	Glu 420	Pro	Ala	Gly	Ser	Asn 425	Thr	Ser	Ser	Phe	Met 430	Pro	Thr
G]	.u	Gly	Thr 435	Ala	Gly	Ser	Leu	Ser 440	Pro	Leu	Pro	Asp	Asp 445	Thr	Ala	Thr
Le	u	Val 450	Leu	Glu	Lys	Arg	Gln 455	Ala	Pro	Leu	Asp	Cys 460	Val	Leu	Tyr	Arg
T)		Gly	Ser	Phe	Ser	Leu 470	Thr	Leu	Asp	Ile	Val 475	Gln	Gly	Ile	Glu	Ser 480
Al	.a	Glu	Ile	Leu	Gln 485	Ala	Val	Ser	Ser	Ser 490	Glu	Gly	Asp	Ala	Phe 495	Glu
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aacaagtgtg aatttggggt tgcacacctg ctctggtttt tctctcccta aaatggaaga
                                                                    180
                                                                    240
tatcaqtaqt qcttcaqqtq tctcccaccc atttqattta qtqaqqacat qqqcaactqa
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                                                                    120
                                                                    180
aacaagtgtg aatttggggt tgcacacctg ctctggtttt tctctcccta aaatggaaga
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cagtagtgct tcaggtgtct cccacccatt tgatttagtg aggacatggg caactgagct
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